

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

08/973303

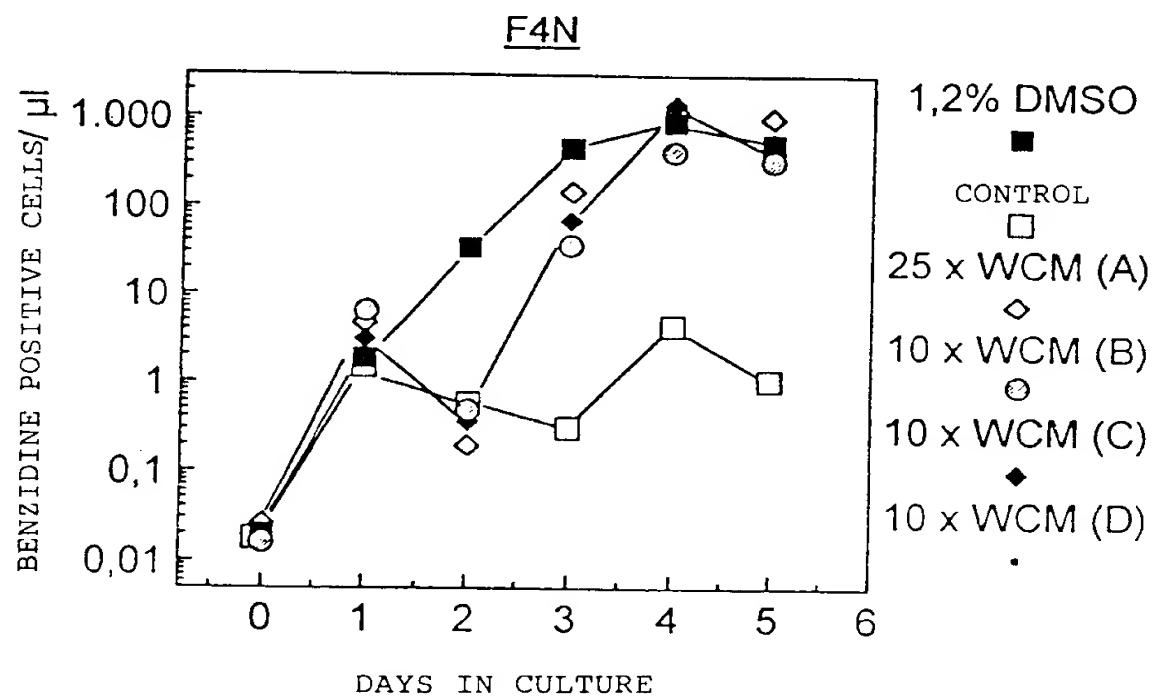


FIGURE 1

APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

08/973303

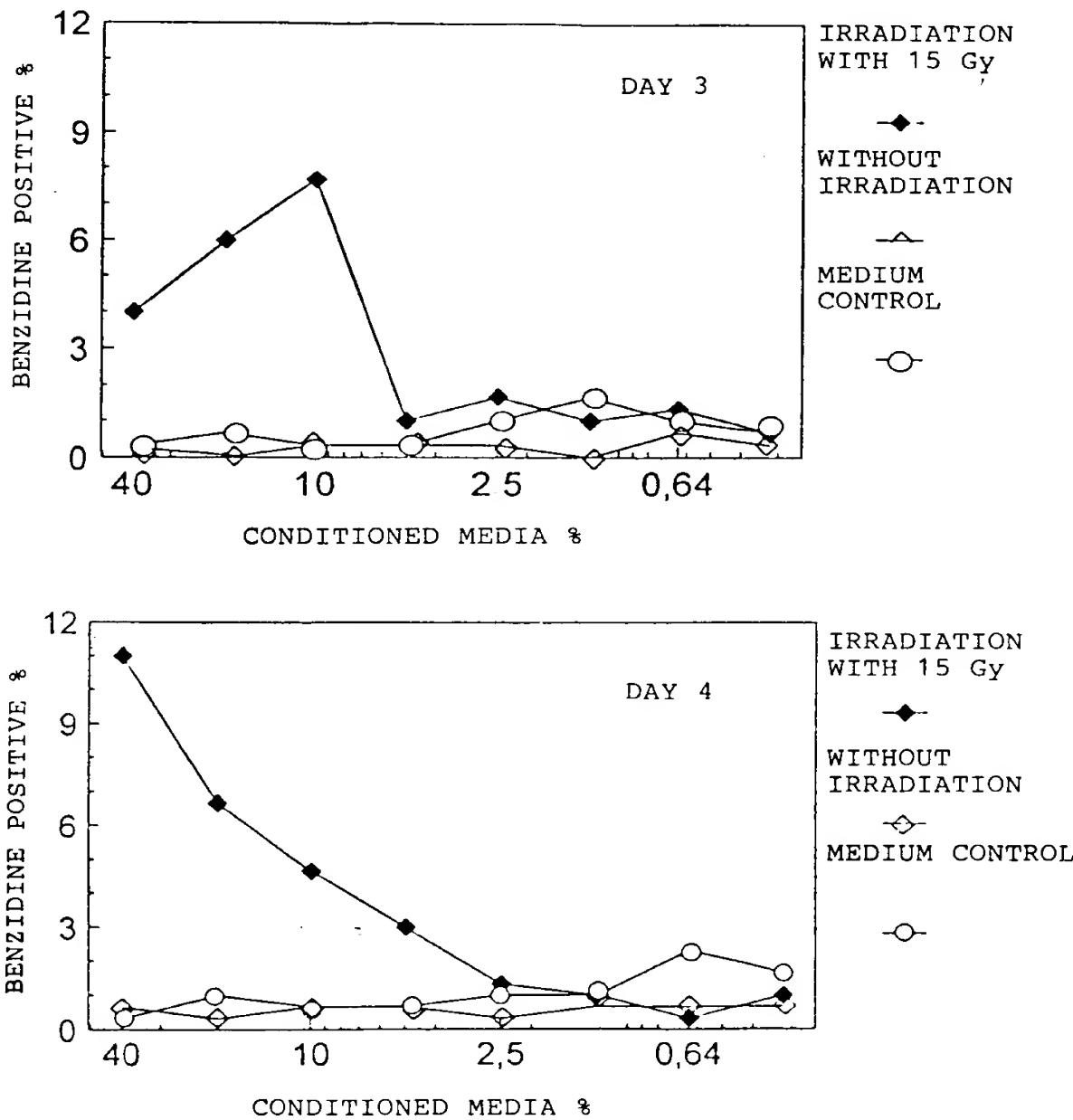


FIGURE 2

APPROVED BY	O.G. FIG.	
DRAFTSMAN	CLASS	SUBCLASS

08/973303

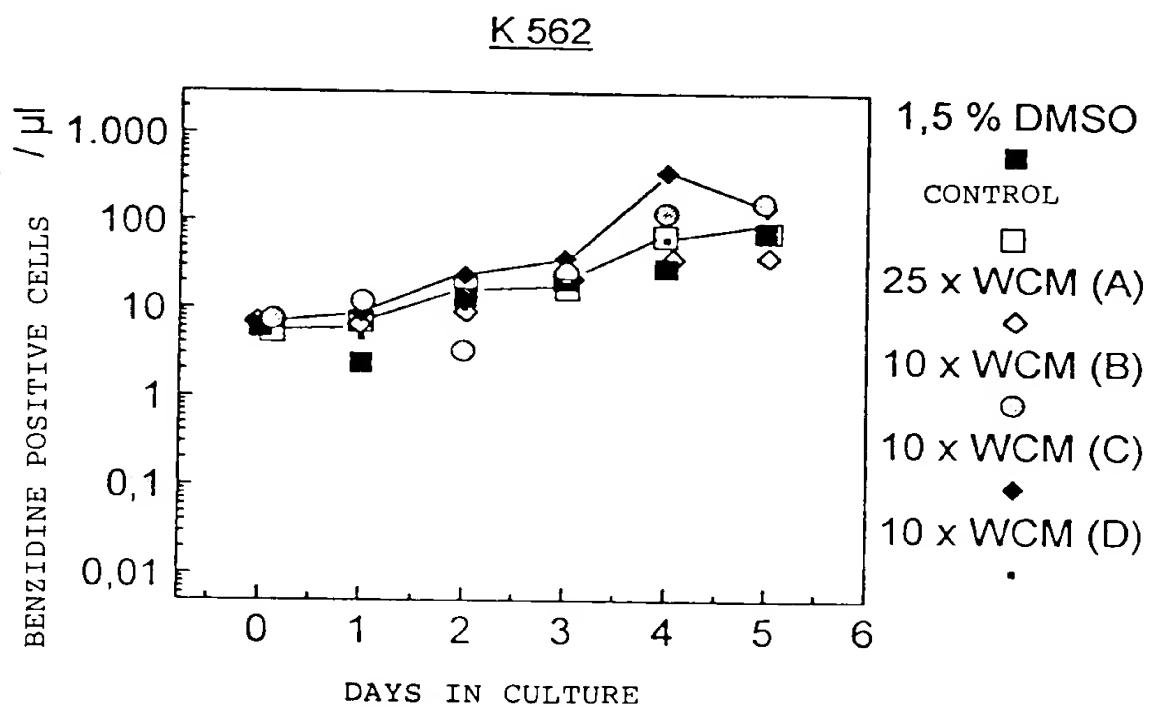


FIGURE 3

APPROVED BY	O.G. FIG.	
DRAFTSMAN	CLASS	SUBCLASS

08/973303

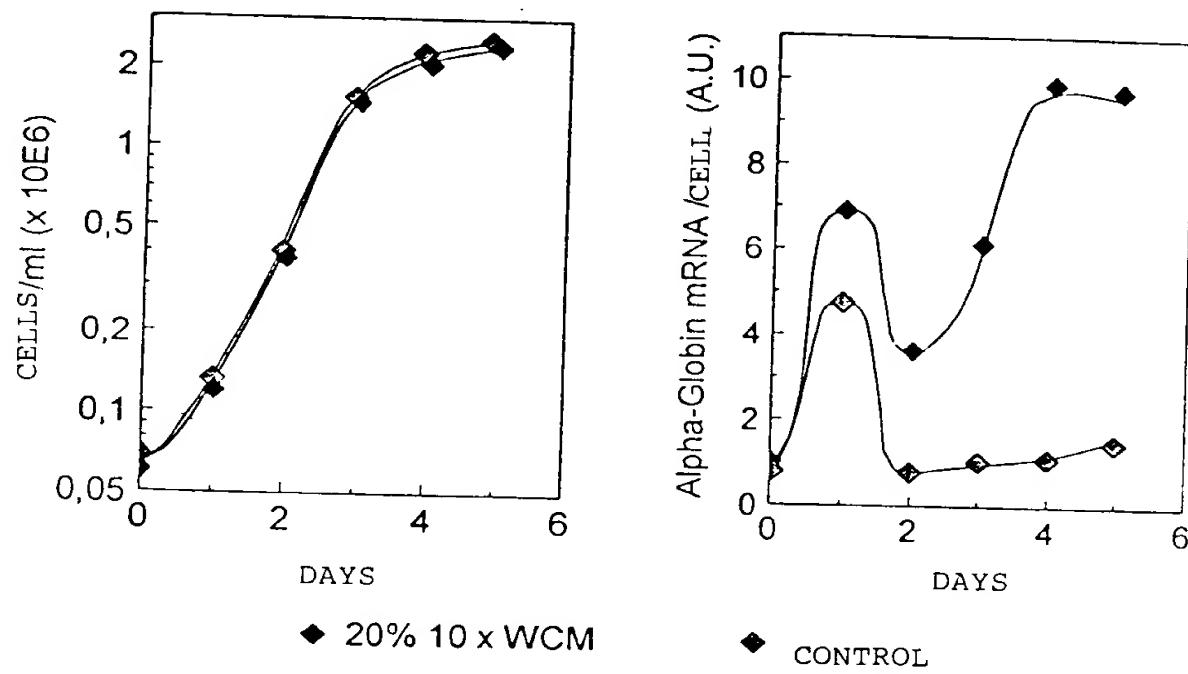


FIGURE 4

APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

08/973303

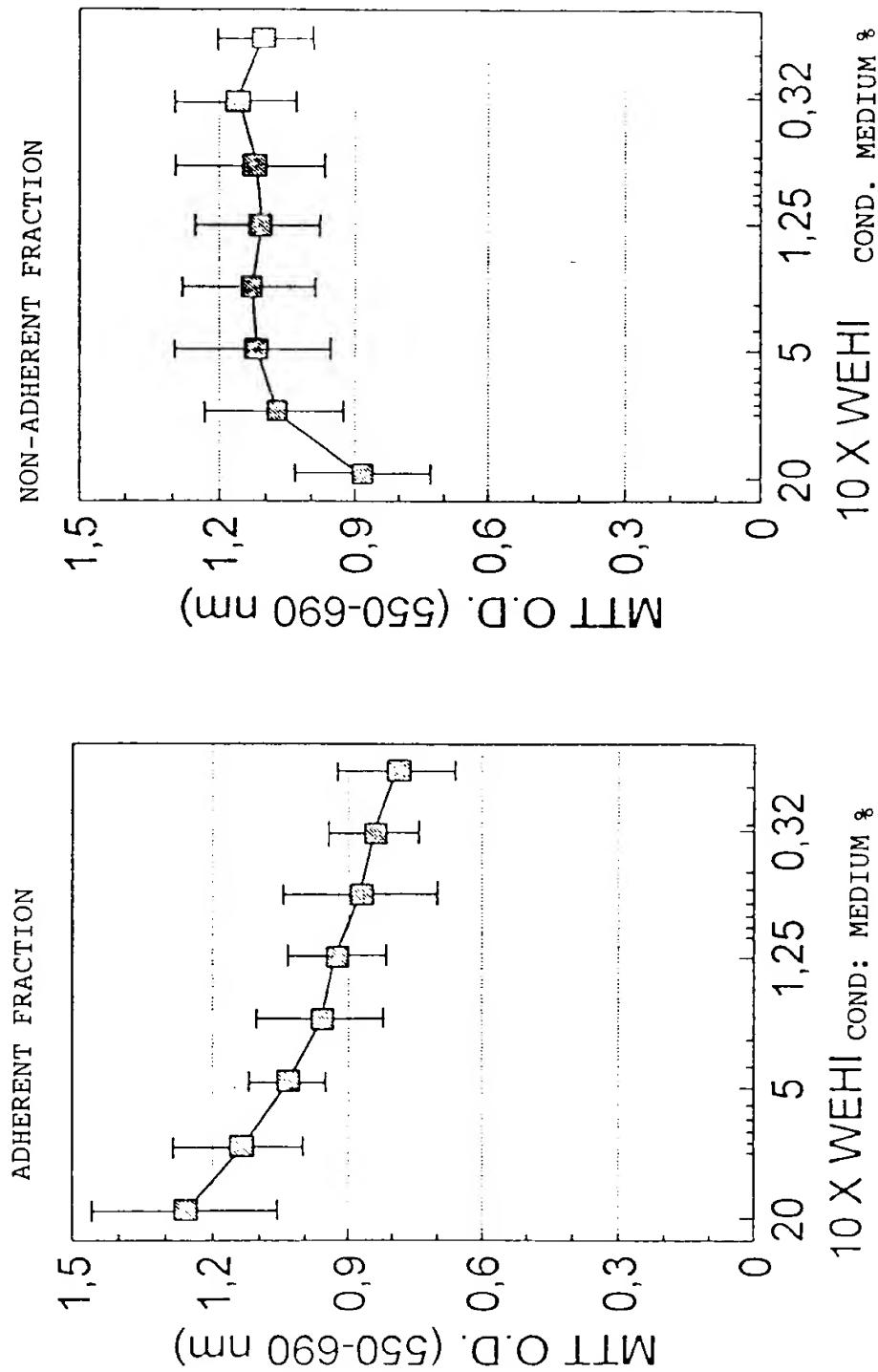


FIGURE 5

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

08/973303

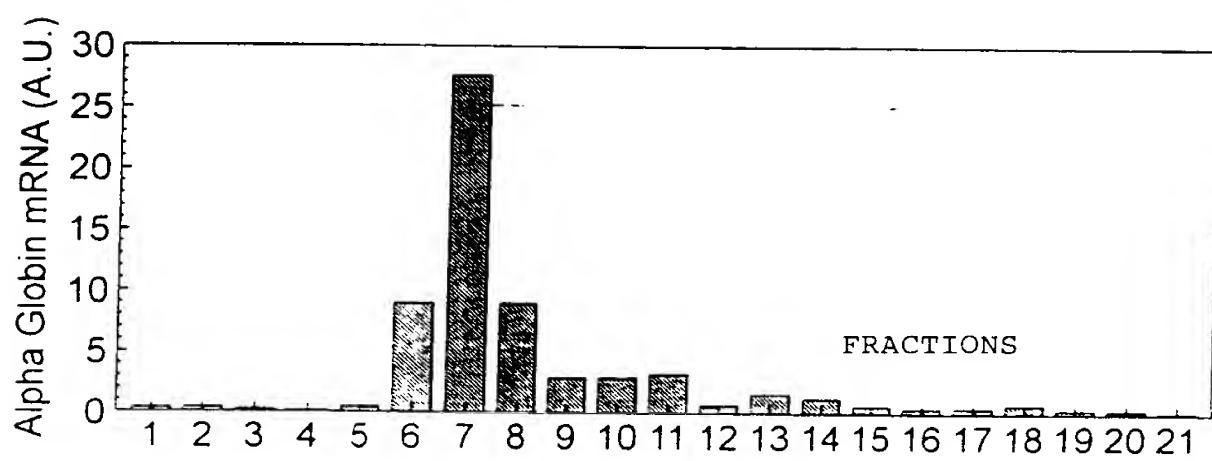


FIGURE 6

APPROVED	O.G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

08/973303

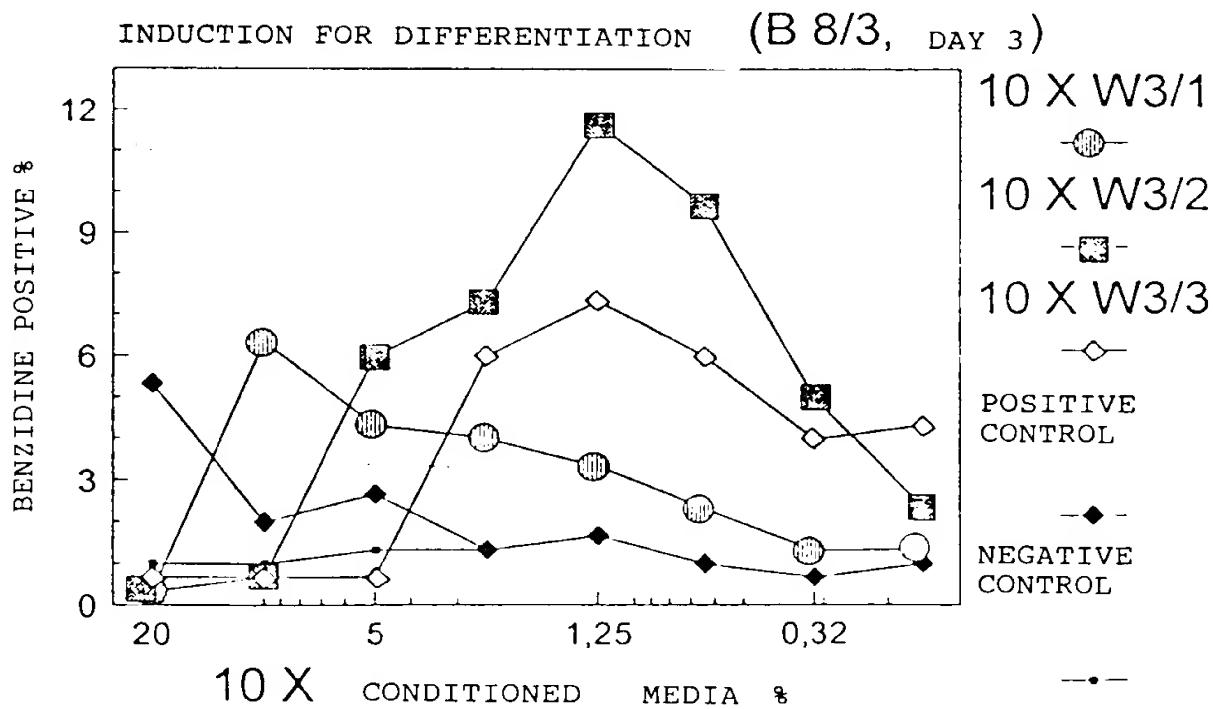
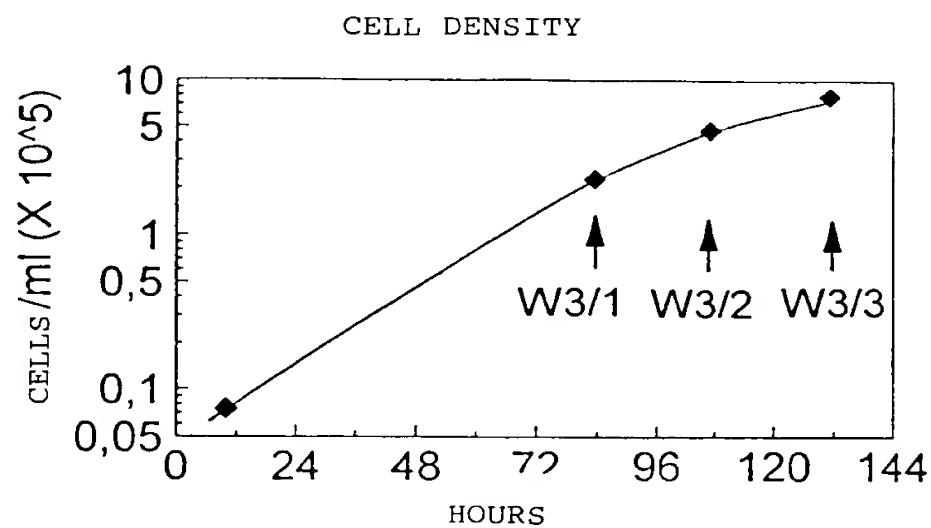
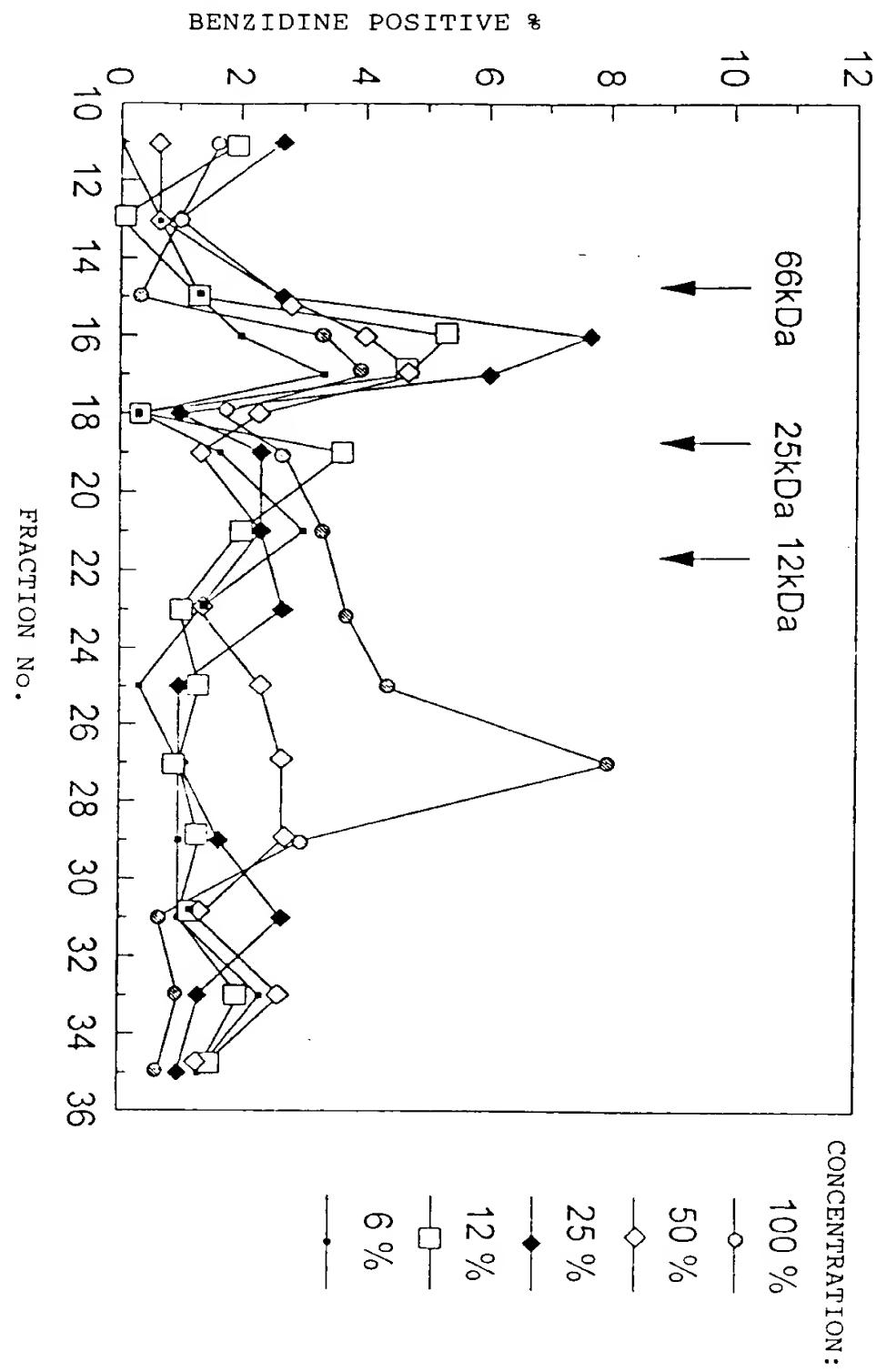


FIGURE 7

08/973303

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS	SUBCLASS
-----------------------------	--------------------	----------



APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

U8/973303

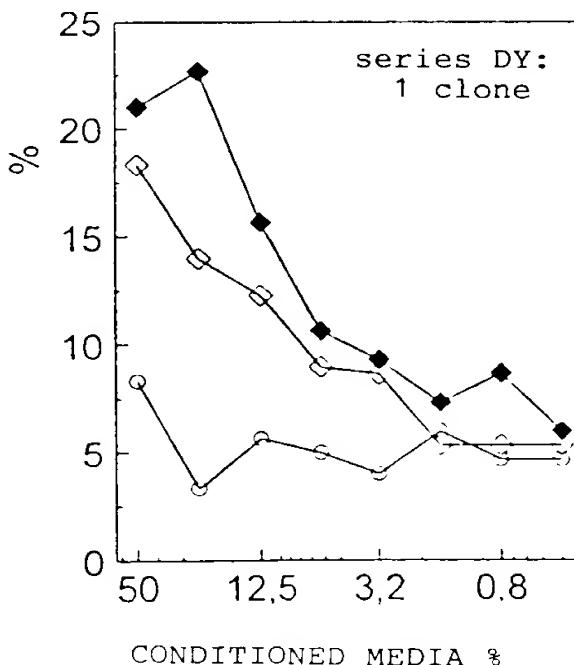
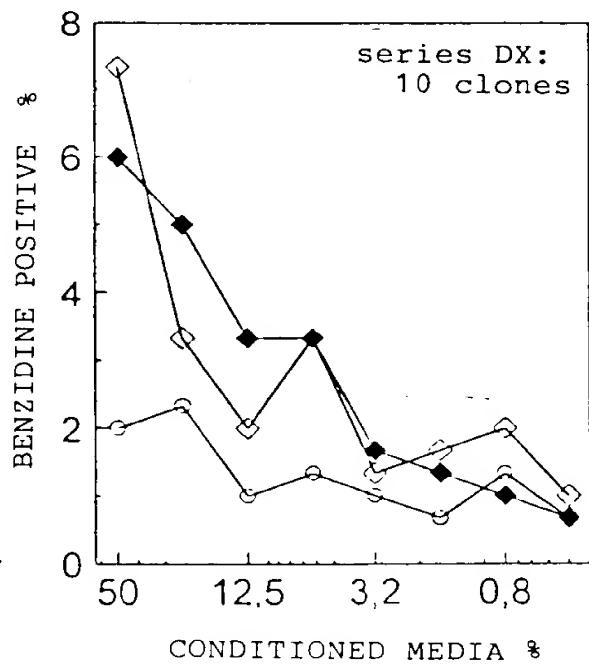
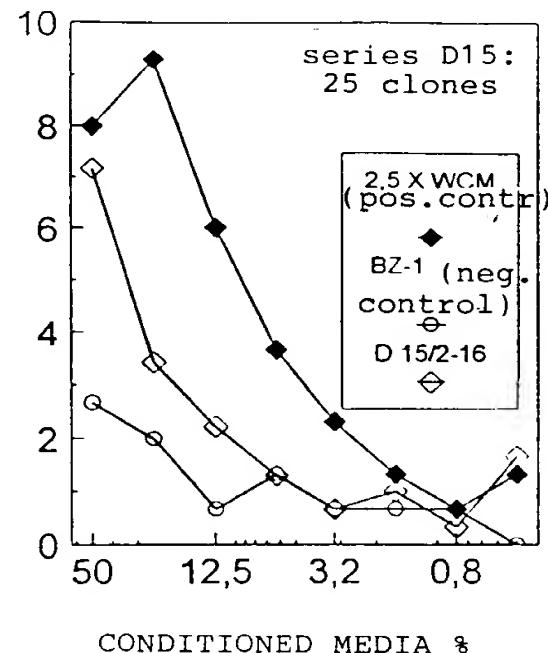
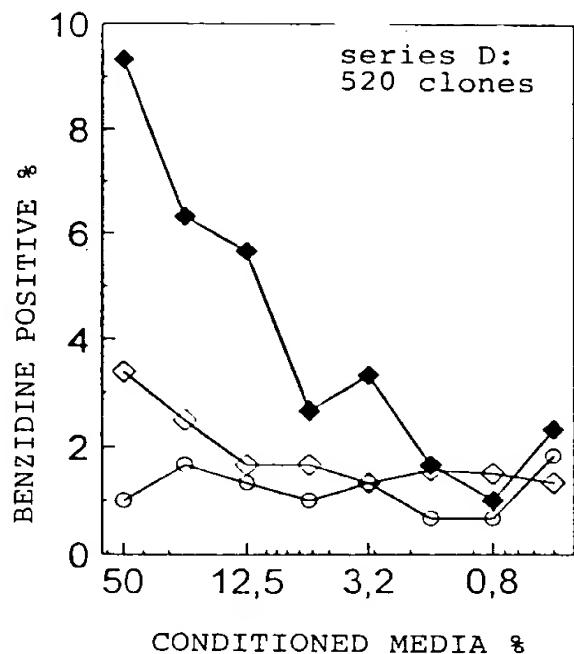


FIGURE 9

APPROVED BY	O.G. FIG.	
	CLASS	SUBCLASS
DRAFTSMAN		

08/973303

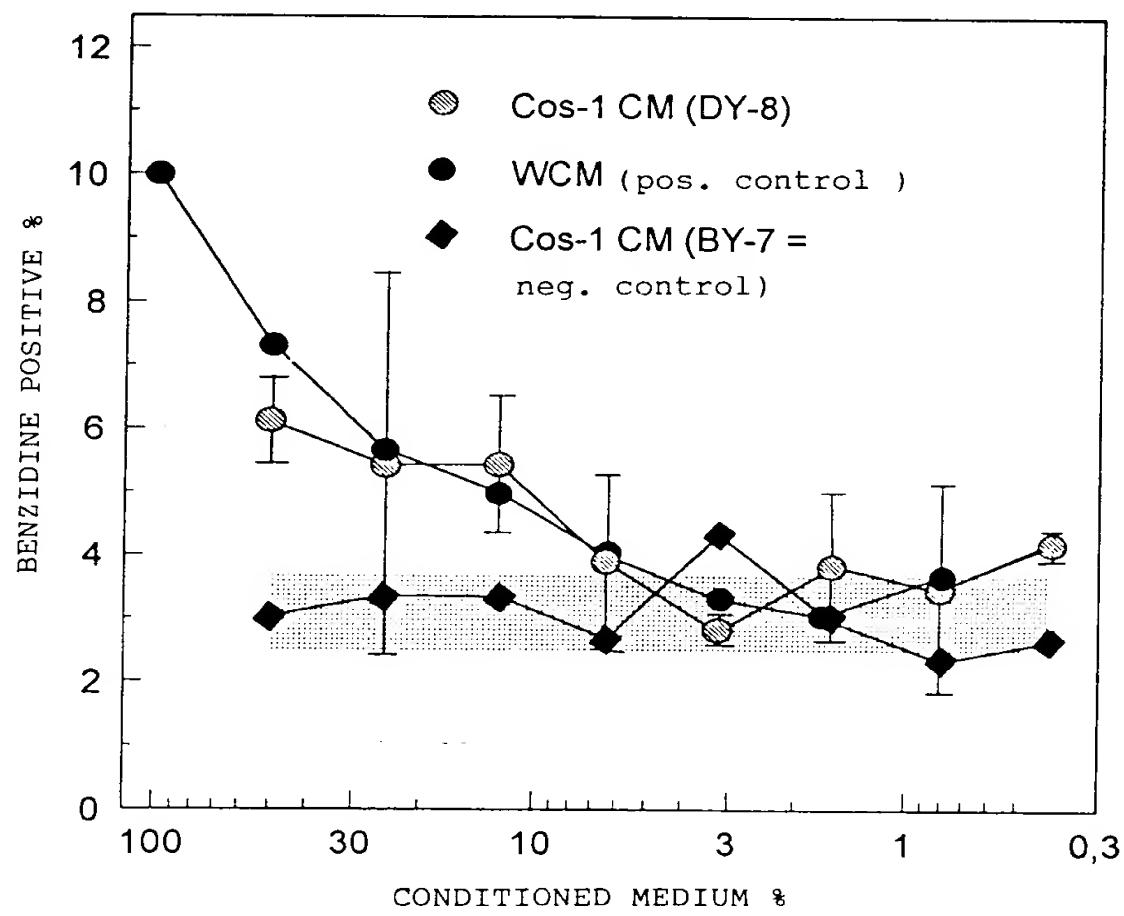


FIGURE 10

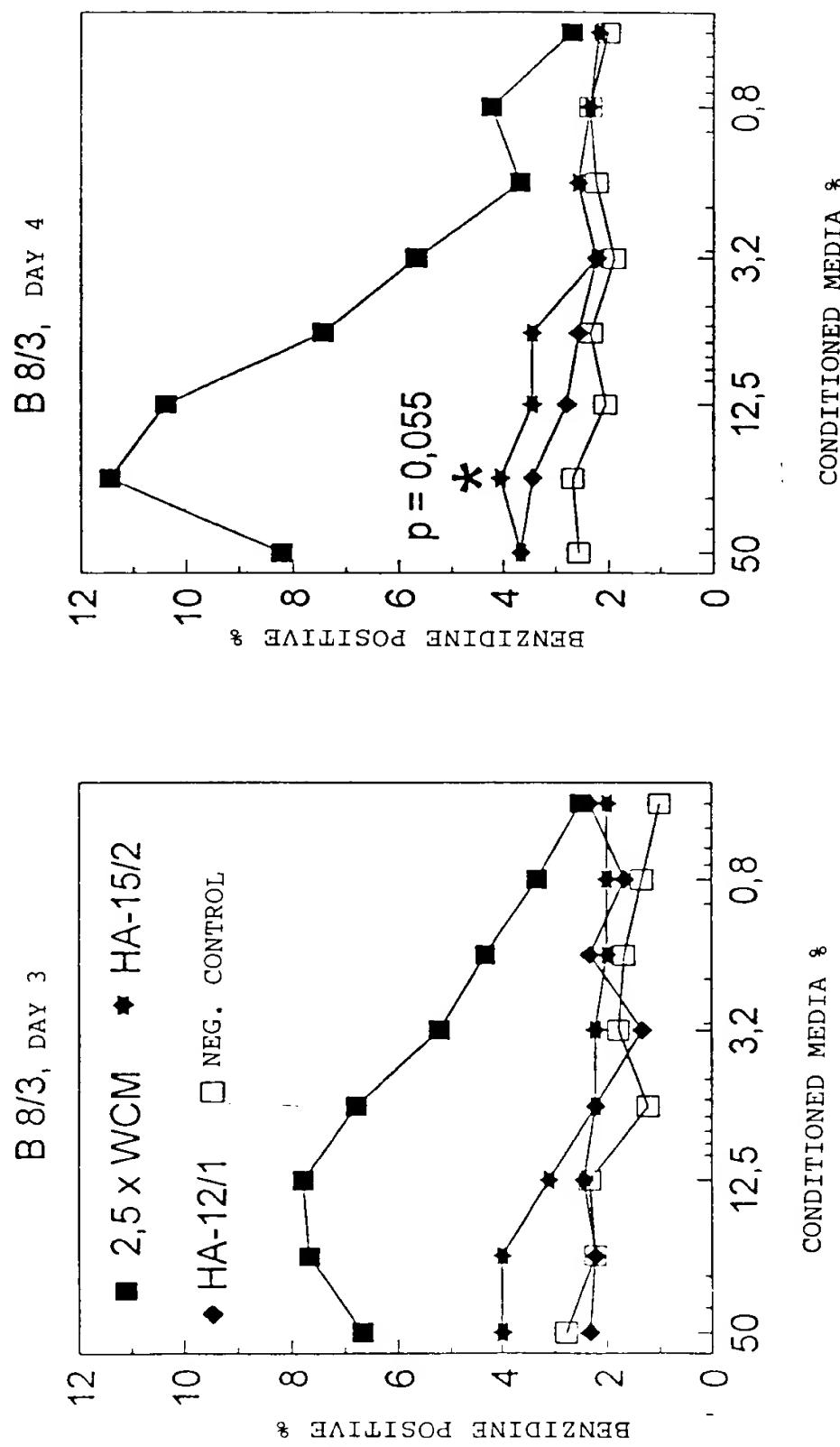


FIGURE 11

APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

08/973303

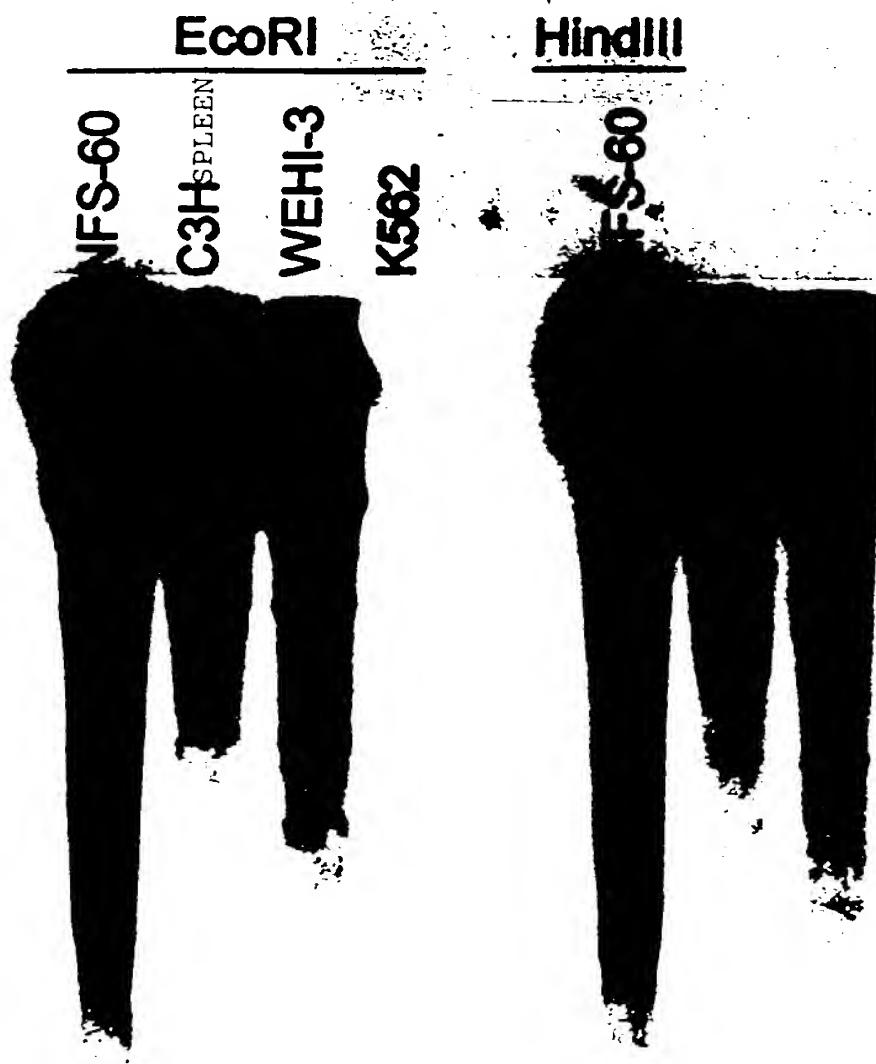
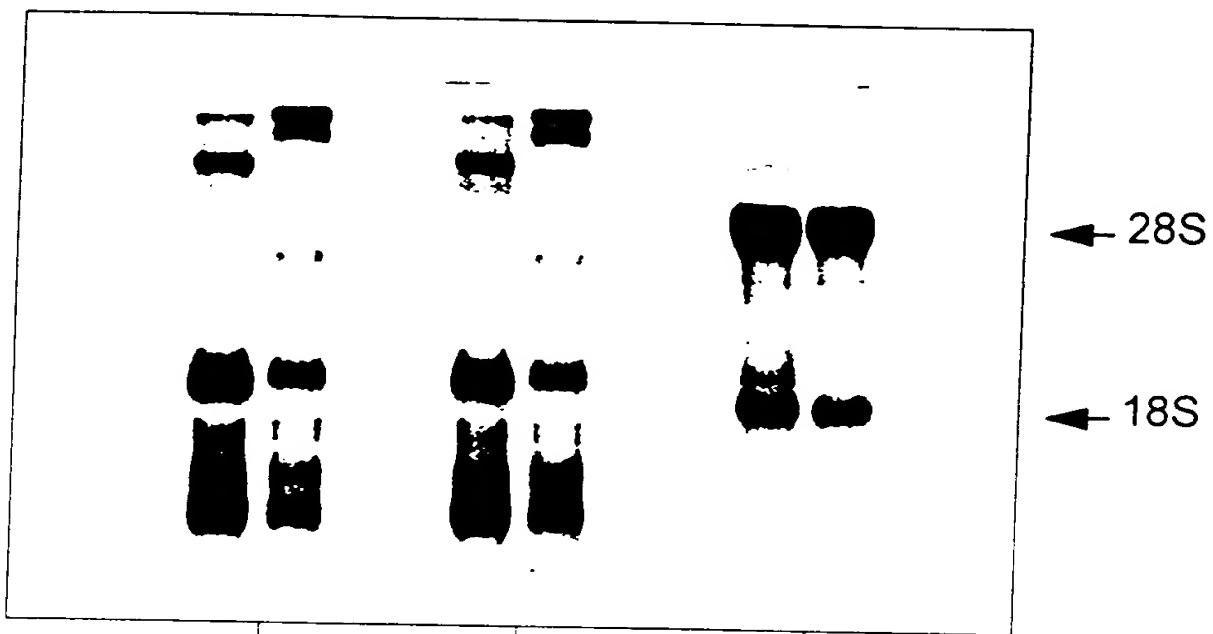


FIGURE 12

08/973303

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		



SAMPLE: 1350 bp 500 bp 3' 220 bp 5'

FIGURE 13

APPROVED	O.G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

08/973303

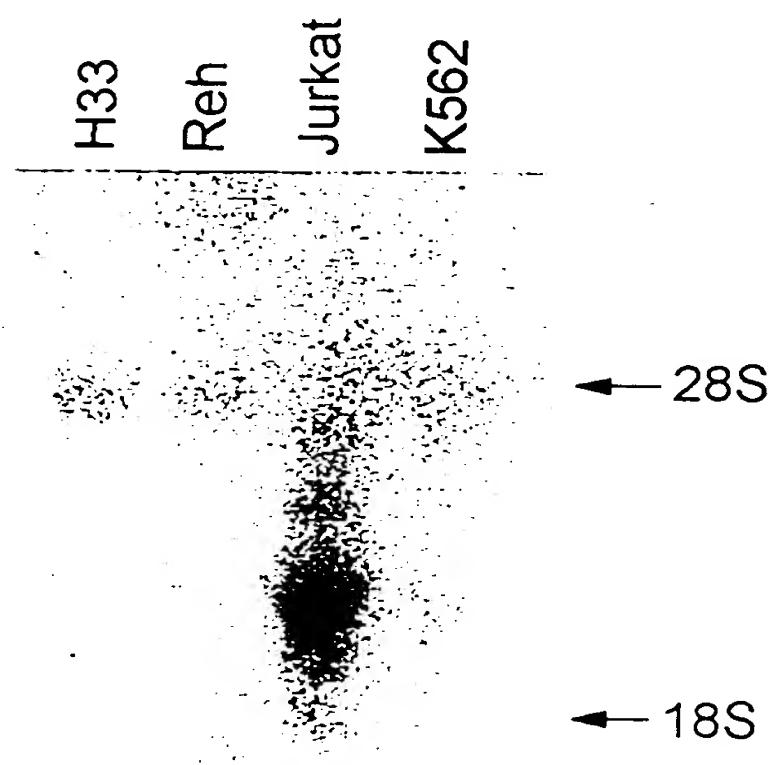


FIGURE 14

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS SUBCLASS
-----------------------------	-------------------------------

08/973303

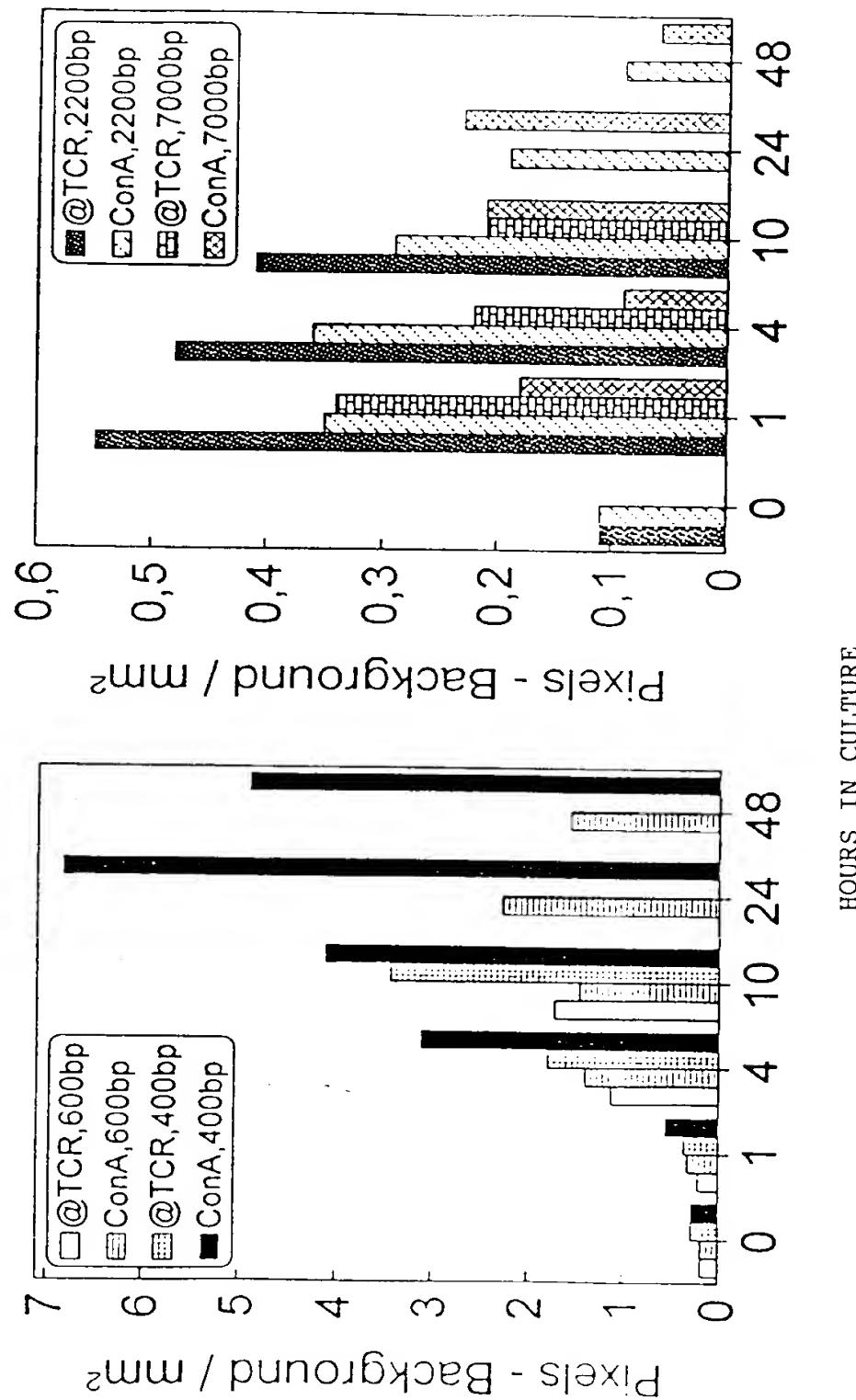


FIGURE 15

08/973303

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

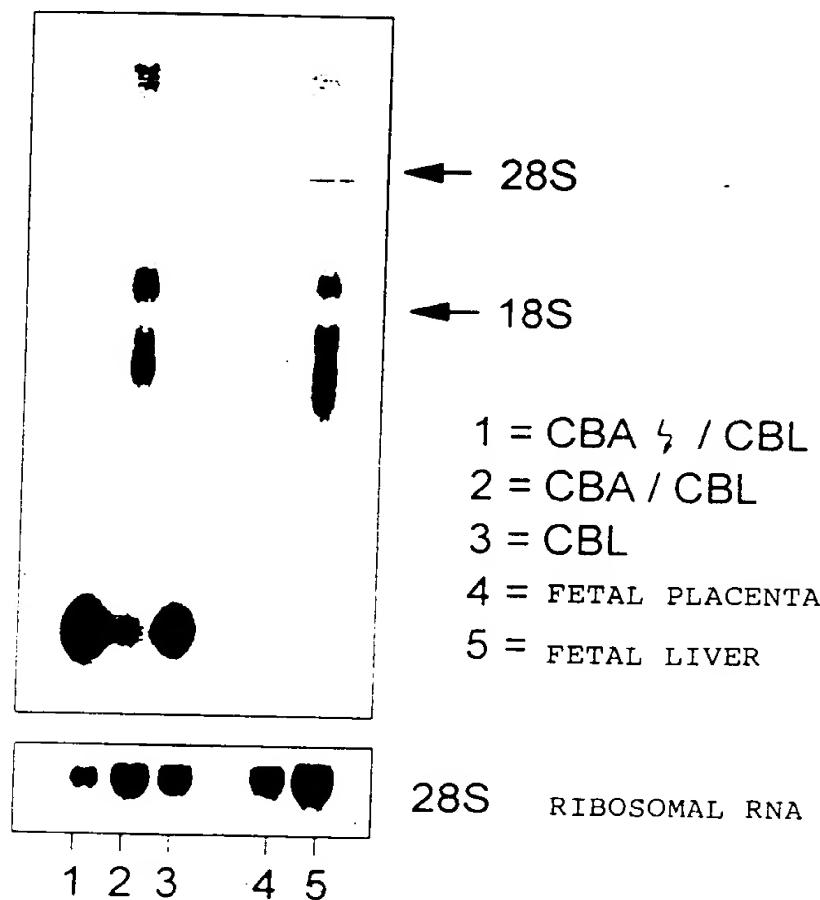


FIGURE 16

APPROVED BY	O.G. FIG.	
DRAFTSMAN	CLASS	SUBCLASS

08/973303

TRANSPLANTATION OF CBA X CBL
OR CBL SPLEEN CELLS

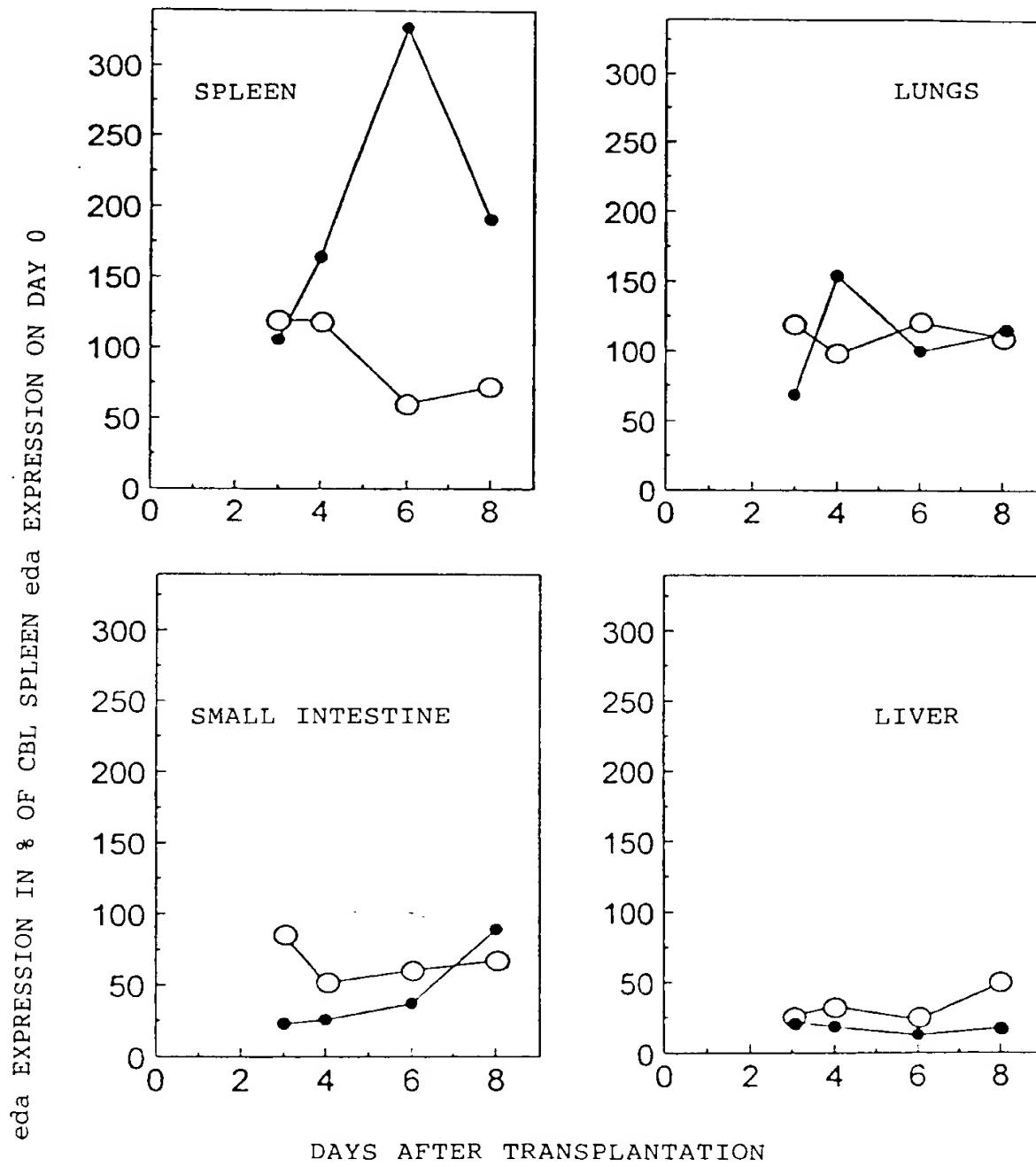


FIGURE 17

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS	SUBCLASS
-----------------------------	--------------------	----------

08/973303

1 CCGACCGTGC GGACTTAAGA TGGAGGCAGT TCCTGTCTGC GGCGGGAAAGA
 51 GAAGGCTCGG TCGGAGCCGG GAATGCTGGG ACTTGTAGTG CGTAGTCAAT
 101 GGTTCTCTAT GGGCTTCAG AGTAGTGGC GGGAAAGGCGG CCCCAGGGCA
 151 TGCTGGGAGT TGTAGTCCTG CCGTCGTCAA TGGTTCTCTA TGGGCTTTCA
 201 GAGTGAGTGG CGGGAAAGGCG GCCCCGAGGC ATGCTGGGAG TTGTAGTCCT
 251 GCCATAGTCA ATGGTTCTCT ATGGGCTTTC AGACTGAGTG GCAGGGAAAGGC
 301 GGCCCCGAGG CATGCTGGGA GTTGCAGCGC CATGTTTAA AGCACCGCGTT
 351 TCTCTGTATA GACCTGGCTG TGGATTTTC GCTAATTCTT TTTTTAGCT
 401 TTATTTTAA TTTTACTTT TTCACACAGG ATTTCTCTT ATAGCCTTGG
 451 CTACCGTTT TTCCCTAATT ATTCTCCTT TCATTTGGT TTATTTTTT
 501 TTAATTTGG TTTTTTAAG ACAGGGTTTC TCTGTATAGA CCTGGCTGTG
 551 GATTCTCAC TAATTATTT TTTAGCTT ATTTTAATT TTTACTTTT
 601 CACACAGGAT TTCTCTTTAT AGCCTGGCT ACCGTTTTT CCGTAATTAT
 651 TCTTATTTTC ATTTGGTTT ATTTTTAAT TTTAATTTTT GATTTGGAG
 701 ACAGGGTTTC TCTTTAGCC GCAGCTATGG TTTCTGCCCT AATTATTCTT
 751 GTCCTTATTT GTAATTAAAT TCTTAATTAA ATTAAATTAA TAATTTGTT
 801 GTAAGTTTT CTGTGGCGT GAATGGAAAG TCTAACCGT GTTCTCTGT
 851 TCAGCGTCCG CCGGTACCGG CCGCCGCC CAGCGACGTC ACCCACACGC
 901 GCAGAAGCGG ACGCCGCGGT CAAGATGTCT CTGCCATGCC CACGGGACGC
 951 ACGGACGCAC GGACGGACGG ACGGACTCCA CAAGGTAGGA AGCCTGCGCC
 1001 GACCGCACCG CCGCACCCAC CACAGCACAC AGGACACACG CGGGCCCCGC
 1051 GCGCGCCAG GCACACCGGG CACACACGGC ACACACGGCA GGCAGGCCAG
 1101 GCACACGCAT CCGCAGGACC CGCGCACCC GCCACGCAGA CACGGACGAG
 1151 CCGCCGCGGT CAAGATGTT ACCCGCCCGC GTCAAGATGT ATGTGCCACC
 1201 GACCCCTCGCC CCGCTGGACG GACGGACGGA CGCACGCACG CGTCAGCGT
 1251 CCACCGGTCA CTGCCGCCGC CCACAGTGAT GTCACCCACG AAAGCACACA
 1301 CGTAGAAGCG GACGCCGTGG TCAAGATGTC TCTGCCATCC CCACAGGACG
 1351 GACGGACGGA CTCCACAAGG TGCCTGTGTC GCCGAGGCCG CCAGGACGGA
 1401 GCGATTCTCA CGGAGGAAGG AGCACGCCAA CAGGGCCTGA CTGCGTACAG
 1451 ACATGTCCCC CTCATAAAA TTGCAGTTGA AATGGAAAAA AAAAAA

FIGURE 18

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

08/973303

PRELIMINARY STRUCTURE OF eda 2.2 kbp cDNA

* = stopcodon, R = Repeat

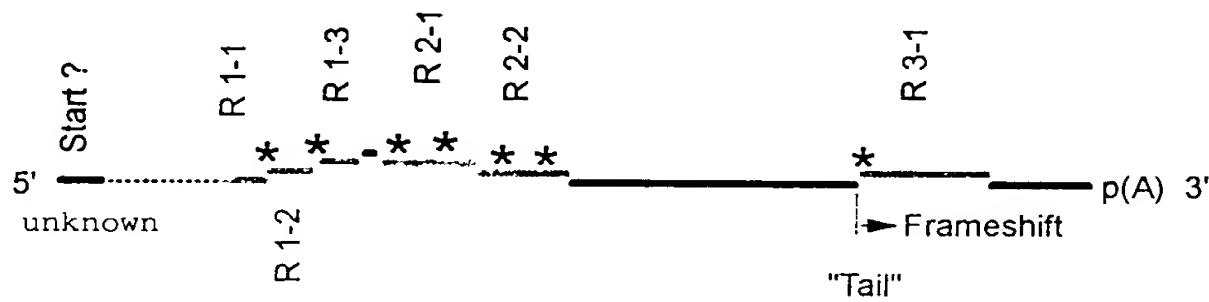


FIGURE 19

APPROVED	O.G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

08/973303

1 CGCGCCCGCC CGGGATCCCC AGCTGCCGCC GCGCCCGCCC GCCCGCCCGG
CONSENSUS SEQUENCE
51 GGCCCCCGCT GCAGAACCGT GACCGTCCGC CGGTACGGC CGCCGCCCGC
ATGGGGCT GCAGAACCGT GACCGTCCGC CGGTACGGC CGCCGCCCGC
101 AGCGACGTCA CCCACACGCG CAGAACGGGA CGCCGCCGGTC AAGATGTCTC
AGCGACGTCA CCCACACGCG CAGAACGGGA CGCCGCCGGTC AAGATGTCTC
151 TGCCATGCCC ACAGGGACGCA CGGACGCACG GACGGACGGGA CTGACTCCAC
TGCCATGCC ACAGGGACGCA CGGACGCACG GACGGACGGGA CTGACTCCAC
201 AAGGTAGGAA GCCTGCGCCG ACCGCACCGC CGCACCCACC ACAGCACACA
AAGGTAGGAA GCCTGCGCCG ACCGCACCGC CGCACCCACC ACAGCACACA
251 GGACACACGC GGGCCCCGCG CCCGCCAGG CACACGGC ACACACGGCA
GGACACACGC GGGCCCCGCG CCCGCCAGG CACACGGC ACACACGGCA
301 CACACGGCAG GCAGGCCAGG CACACGCATC CGCAGGACCC GCCGCACCCG
CACACGGCAG GCAGGCCAGG CACACGCATC CGCAGGACCC GCCGCACCCG
351 CCACGCAGAC ACGGACGAGC CGCCGCCGGTC AAGATGTTCA CCCGCCGG
CCACGCAGAC ACGGACGAGC CGCCGCCGGTC AAGATGTTCA CCCGCCGG
401 TCAAGATGTA TGTGCCACCG ACCCTCGCCC CGCTGGACGG ACGGACGGAC
TCAAGATGTA TGTGCCACCG ACCCTCGCCC CGCTGGACGG ACGGACGGAC
451 GCGGCCACGC CGTCAGCGTC CACCGGTAC TGCCGCCGCC CACAGTGACG
GCGGCCACGC CGTCAGCGTC CACCGGTAC TGCCGCCGCC CACAGTGACG
501 TCACCCACGA AAGCACACAC GTAGAACGGG ACGCCGTGGT CAAGATGTCT
TCACCCACGA AAGCACACAC GTAGAACGGG ACGCCGTGGT CAAGATGTCT
551 CTGCCATCCC CACAGGACGG ACGGACGGAC TCCACAAGGT GCGCGTGTG
CTGCCATCCC CACAGGACGG ACGGACGGAC TCCACAAGGT GCGCGTGTG
601 CCGAGGCCGC CAGGATGGAG CGATTCTCAC GGAGGAAGGA GCACGCCAAC
CCGAGGCCGC CAGGATGGAG CGATTCTCAC GGAGGAAGGA GCACGCCAAC
651 AGGGCCTGAC TGCCTACAGA AATGCCCGCC CTCAATAAAA TTGCAGTTGA
AGGGCCTGAC TGCCTACAGA AATGCCCGCC CTCAATAAA
701 AATGGAAAAA AAAAAA

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS SUBCLASS
-----------------------------	----------------------------------

08/973303

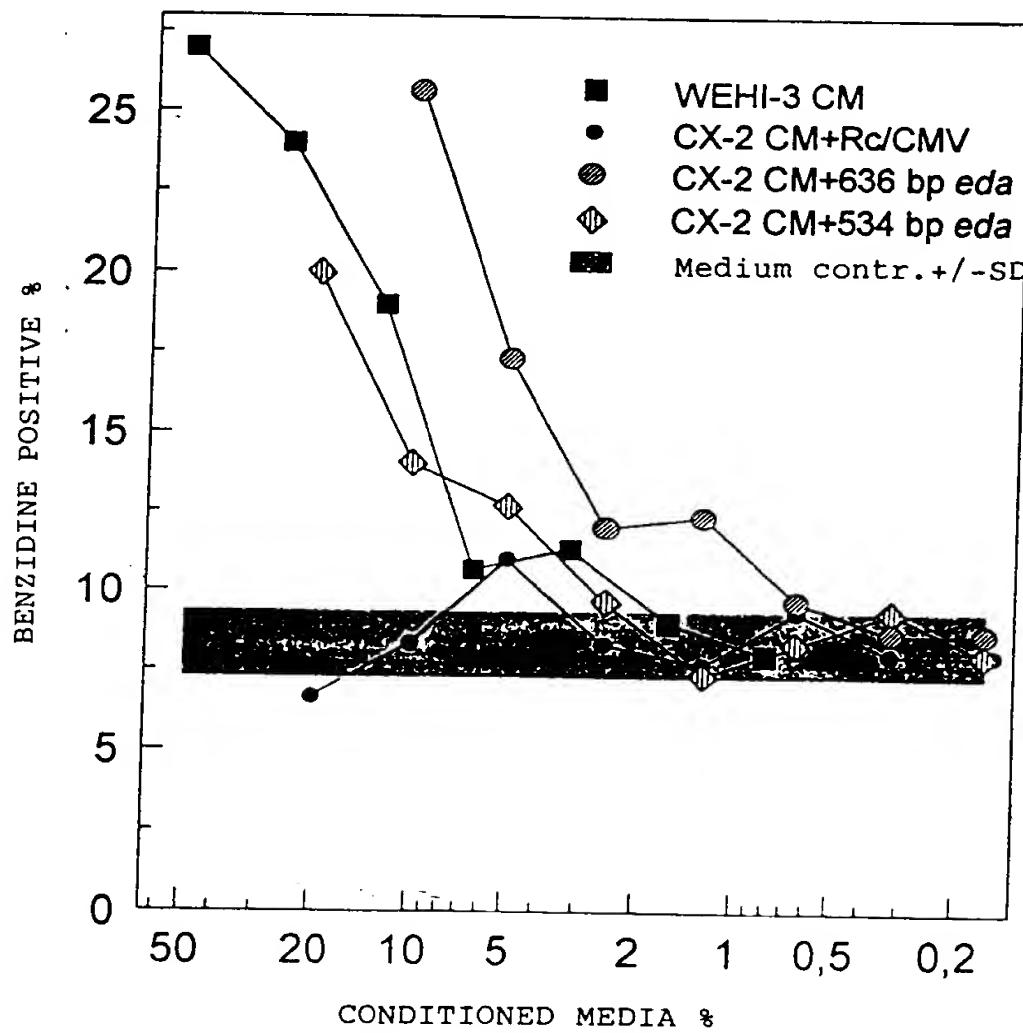
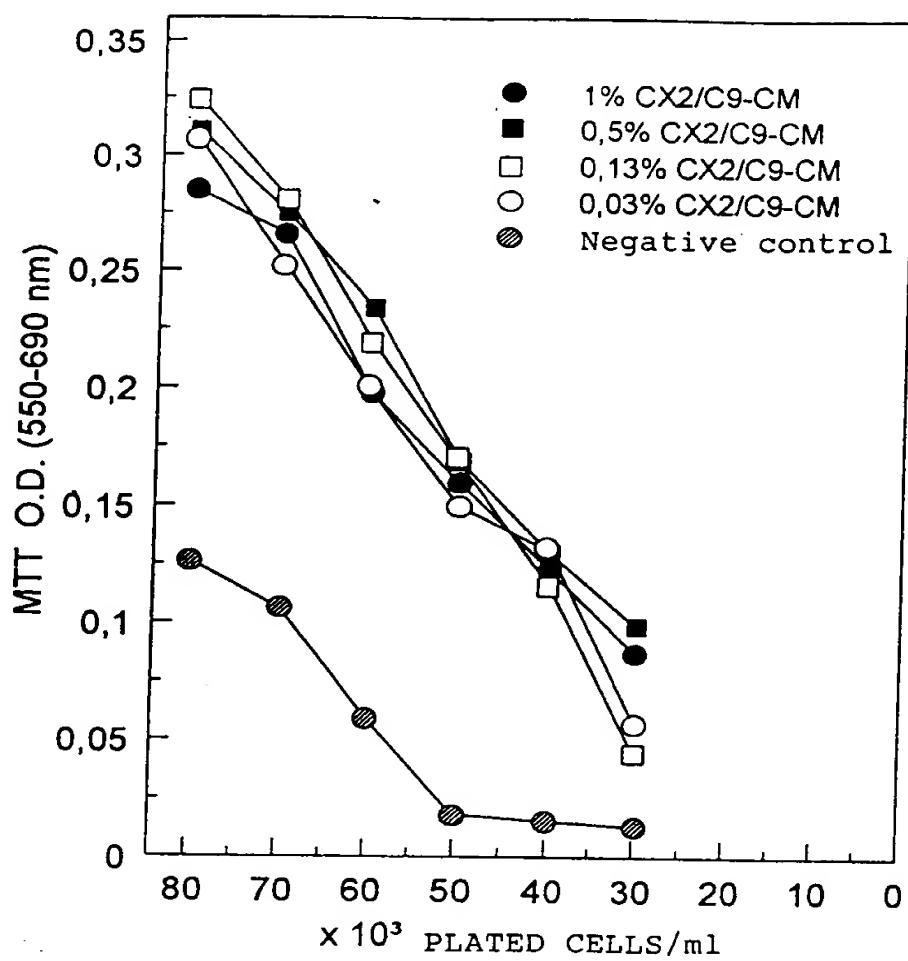


FIGURE 21

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS SUBCLASS
-----------------------------	----------------------------------

08/973303

A



B

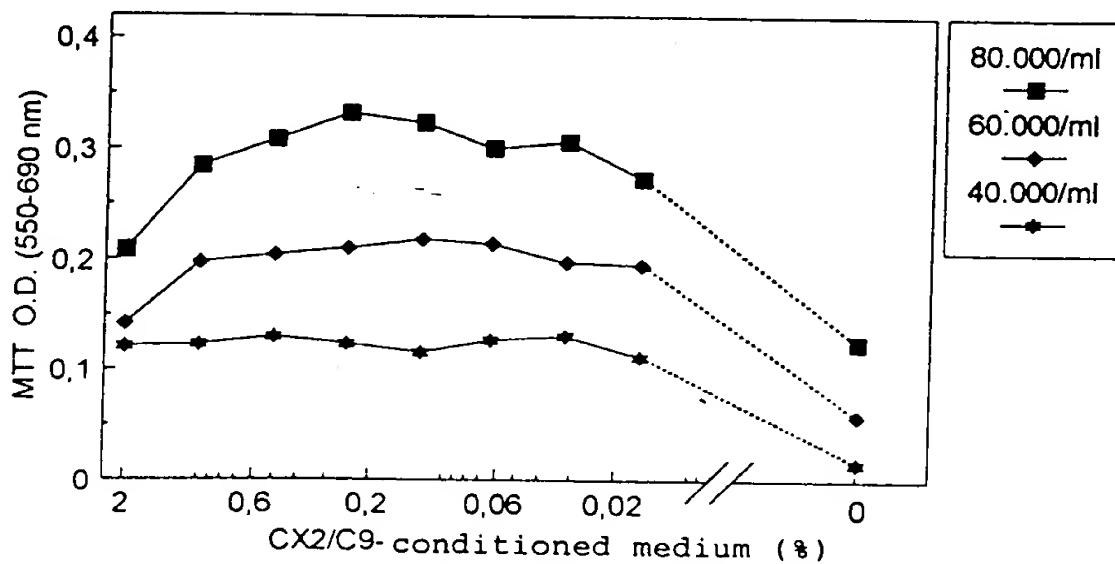


FIGURE 22

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS	SUBCLASS
-----------------------------	--------------------	----------

08/973303

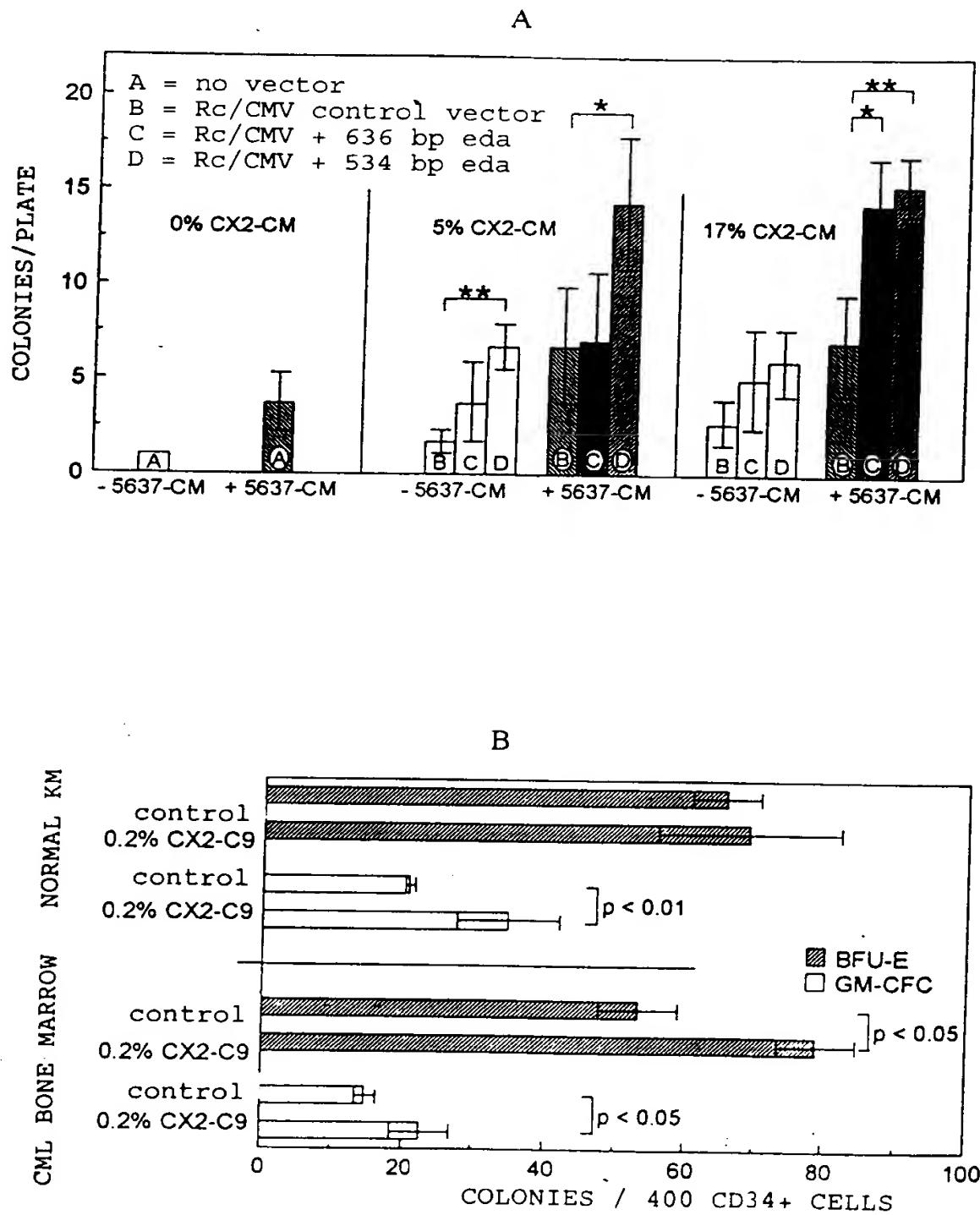


FIGURE 23